Amendments to the Claims

We claim:

- 1. (Currently amended) An <u>isolated recombinant purified</u> CA125 molecule, comprising:
- (a) an extracellular amino terminal domain, comprising amino acids #1-1637 #1-33 of SEQ ID NO: 299, amino acids #34-1593 of SEQ ID NO: 299, amino acids #1594-1605 of SEQ ID NO: 299, amino acids #1606-1617 of SEQ ID NO: 299, and amino acids #1618-1637 of SEQ ID NO: 299;
- (b) a multiple repeat domain <u>comprising residues 3200 to 3355 of SEQ ID NO: 162 SEQ ID NO:150</u>; and
- (c) a carboxy terminal domain comprising a transmembrane anchor with a short eytoplasmic domain, and further comprising amino acids #1-284 #1-11 of SEQ ID NO: 300; amino acids #12-33 of SEQ ID NO: 300; amino acids #34-82 of SEQ ID NO: 300; amino acids #83-133 of SEQ ID NO: 300; amino acids #134-156 of SEQ ID NO: 300; amino acids #157-212 of SEQ ID NO: 300; amino acids #213-225 of SEQ ID NO: 300; amino acids #226-253 of SEQ ID NO: 300; and amino acids #254-284 of SEQ ID NO: 300.
- 2. (original) The CA125 molecule according to claim 1, wherein N-glycosylation sites of the amino terminal domain marked (x) in FIG. 8B are encoded at positions #81, #271, #320, #624, #795, #834, #938, and #1,165 in SEQ ID NO: 299.
- 3. (original) The CA125 molecule according to claim 1, wherein the serine and threonine O-glycosylation pattern for the amino terminal domain is marked (o) in SEQ ID NO: 299 in FIG. 8B.
- 4. (canceled).

- 5. (previously presented) The CA125 molecule according to claim 1, wherein the multiple repeat domain comprises 156 amino acid repeat units which comprise epitope binding sites.
- 6. (previously presented) The CA125 molecule according to claim 5, wherein the epitope binding sites are located in the C-enclosure at amino acids #59-79 (marked C-C) in SEQ ID NO: 150 in FIG. 5.
- 7. (previously presented) The CA125 molecule according to claim 5, wherein the 156 amino acid repeat unit comprises O-glycosylation sites at positions #128, #129, #132, #133, #134, #135, #139, #145, #146, #148, #150, #151, and #156 in SEQ ID NO: 150 in FIG. 5.
- 8. (previously presented) The CA125 molecule according to claim 5, wherein the 156 amino acid repeat unit comprises N-glycosylation sites at positions #33 and #49 in SEQ ID NO: 150 in FIG. 5.
- 9. (previously presented) The CA125 molecule according to claim 5, wherein the 156 amino acid repeat unit comprises at least one conserved methionine (designated M) at position #24 in SEQ ID NO: 150 in FIG. 5.
- 10. (Currently amended) The CA125 molecule according to claim 1, wherein <u>a the</u> transmembrane anchor of the carboxy terminal domain is located at positions #230-252 (underlined) in SEQ ID NO: 300 of FIG. 9B.
- 11. (Currently amended) The CA125 molecule according to claim 1, wherein <u>a the</u> cytoplasmic domain of the carboxy terminal domain comprises a highly basic sequence adjacent to the transmembrane anchor at positions #256-260 in SEQ ID NO: 300 of FIG. 9B, serine and threonine phosporylation sites at positions #254, #255, and #276 in SEQ ID NO: 300 in FIG. 9B, and tyrosine phosphorylation sites at positions #264, #273, and #274 in SEQ ID NO: 300 of FIG. 9B.

12-36. (canceled)

37. (withdrawn) The purified CA125 molecule of claim 1, wherein the multiple repeat domain comprises multiple repeat units, wherein each repeat unit comprises amino acids #1-42 in any of SEQ ID NOS: 164 through 194; at amino acids #43-65 in any of SEQ ID NOS: 195 through 221; amino acids #66-123 in any of SEQ ID NOS: 222 through 249; amino acids #124-135 in any of SEQ ID NOS: 250 through 277; and amino acids #136-156 in any of SEQ ID NOS: 278 through 298.